



0370
12/19

RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 09 / 927,811B
Source: O IPE
Date Processed by STIC: 2/12/03

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



Does Not Comply
Corrected Diskette Needed

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/927,811B

DATE: 02/12/2003
TIME: 12:29:54

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF4\02122003\I927811B.raw

Please re-submit in
the English
language.

```

3 <110> APPLICANT: RheinBiotech Gesellschaft fur neue biotechnologische Prozesse
4     und Produkte mbH
6     Romano, Ivano
8     Gellissen, Gerd
10    DeVergilio, Claudio
12 <120> TITLE OF INVENTION: Heat-Inducible Promoter
14 <130> FILE REFERENCE: PCT1106-01966
16 <140> CURRENT APPLICATION NUMBER: 09/927,811B
C--> 18 <141> CURRENT FILING DATE: 2003-02-07
20 <150> PRIOR APPLICATION NUMBER: PCT/EP00/01144
22 <151> PRIOR FILING DATE: 2000-02-11
E--> 24 <160> NUMBER OF SEQ ID NOS: (27) - found 28
26 <170> SOFTWARE: PatentIn version 3.1

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ERRORED SEQUENCES

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601 <210> SEQ ID NO: 28
602 <211> LENGTH: 15
603 <212> TYPE: DNA
604 <213> ORGANISM: Saccharomyces cerevisiae
W--> 605 <220> FEATURE:
606 <223> OTHER INFORMATION: Heat shock element
608 <400> SEQUENCE: 28
609 ggaacagaac aatcg
                                         15
W--> 611 (continued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(con
W--> 612 tinued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continu
W--> 613 ed) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continued) . (
W--> 614 continued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(cont
W--> 615 inued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continue
W--> 616 d) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continued) . (c
W--> 617 ontinued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(conti
W--> 618 nued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continued
W--> 619 ) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continued) . (co
W--> 620 ntinued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continued)

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delete

{contin
W--> 621 ued) . (continued) . (continued) . (continued) . (continued) . (continued) .
(continued)
W--> 622 . (continued) . (continued) . (continued) . (continued) .
W--> 623 (8)
E--> 625 (1) -delete

} delete

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 02/12/2003
PATENT APPLICATION: US/09/927,811B TIME: 12:29:56

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF4\02122003\I927811B.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:28; Line(s) 611,612,613,614,615,616,617,618,619,620,621

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/927,811B

DATE: 02/12/2003

TIME: 12:29:56

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\02122003\I927811B.raw

L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:71 M:283 W: Missing Blank Line separator, <220> field identifier
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:86 M:283 W: Missing Blank Line separator, <220> field identifier
L:92 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:109 M:283 W: Missing Blank Line separator, <220> field identifier
L:385 M:283 W: Missing Blank Line separator, <220> field identifier
L:389 M:283 W: Missing Blank Line separator, <220> field identifier
L:395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:402 M:283 W: Missing Blank Line separator, <220> field identifier
L:404 M:283 W: Missing Blank Line separator, <220> field identifier
L:416 M:283 W: Missing Blank Line separator, <220> field identifier
L:430 M:283 W: Missing Blank Line separator, <400> field identifier
L:440 M:283 W: Missing Blank Line separator, <400> field identifier
L:447 M:283 W: Missing Blank Line separator, <220> field identifier
L:484 M:283 W: Missing Blank Line separator, <400> field identifier
L:494 M:283 W: Missing Blank Line separator, <400> field identifier
L:552 M:283 W: Missing Blank Line separator, <400> field identifier
L:563 M:283 W: Missing Blank Line separator, <400> field identifier
L:585 M:283 W: Missing Blank Line separator, <400> field identifier
L:605 M:283 W: Missing Blank Line separator, <220> field identifier
L:611 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:612 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:612 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:613 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
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L:619 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:619 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:620 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:620 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:621 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:621 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:622 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:622 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:623 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
L:625 M:254 E: No. of Bases conflict, this line has no nucleotides.
L:24 M:203 E: No. of Seq. differs, <160> Number Of Sequences: Input (27) Counted (28)